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- 5 26. The protein of claim 1, wherein the variant protein has a Group 4 amino acid residue mutated to a Group 5 amino acid residue at position 153.
 - 27. The protein of claim 26 having the mutation C153R.

28. The protein of claim 1, wherein the variant protein has a Group 4 amino acid residue mutated to a Group 1

15 29. The protein of claim 28 having the mutation T281A.

amino acid residue at position 281.

30. The protein of claim 1, wherein the variant protein has Group 3 amino acid residue mutated to a Group 2 amino acid residue at position 367.

31. The protein of claim 30 having the mutation N367I.

- 32. The protein of claim 1, wherein the variant protein has a Group 3 amino acid residue mutated to a Group 6 amino acid residue at position 367.
- 33. The protein of claim 32 having the mutation N367Y.
- 34. The protein of claim 1, wherein the variant protein 30 has a Group 1 amino acid residue mutated to Group 4 amino acid residue at position 389.
 - 35. The protein of claim 34 having the mutation P389S.
- 36. The protein of claim 1, wherein the variant protein has a Group 1 amino acid residue mutated to a Group 2 amino acid residue at position 389.
- 37. The protein of claim 36 having the mutation P389L. 40
 - 38. The protein of claim 1 selected from the group consisting of SEQ ID NO:42, SEQ ID NO:43, SEQ ID NO:44, SEQ ID NO:45, SEQ ID NO:46, SEQ ID NO:47, SEQ ID NO:48,

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- 5 SEQ ID NO:49, SEQ ID NO:50, SEQ ID NO:51, SEQ ID NO:53, SEQ ID NO:54, SEQ ID NO:56, SEQ ID NO:57, SEQ ID NO:58, SEQ ID NO:59, SEQ ID NO:61, SEQ ID NO:62, SEQ ID NO:63, SEQ ID NO:64, and SEQ ID NO:65.
- 39. A nucleic acid comprising a polynucleotide sequence encoding an amino acid sequence of a variant protein of the lovE protein having at least one mutation selected from the group consisting of:
 - (a) a Group 6 amino acid residue mutated to a Group 2 amino acid residue at position 31;
 - (b) a Group 3 amino acid residue mutated to a Group 5 amino acid residue at position 41;
 - (c) a Group 4 amino acid residue mutated to a Group 2 amino acid residue at position 52;
 - (d) a Group 4 amino acid residue mutated to a Group 3 amino acid residue at position 52;
 - (e) a Group 4 amino acid residue mutated to a Group 5 amino acid residue at position 73;
 - (f) a Group 1 amino acid residue mutated to a Group 4 amino acid residue at position 101;
 - (g) a Group 1 amino acid residue mutated to a Group 3 amino acid residue at position 101;
 - (h) a valine amino acid residue mutated to another Group 2 amino acid residue at position 111;
 - (i) a Group 4 amino acid residue mutated to a Group 2 amino acid residue at position 133;
 - (j) an Group 3 amino acid residue mutated to a Group 2 amino acid residue at position 141;
 - (k) an Group 3 amino acid residue mutated to a Group 5 amino acid residue at position 141;
 - (1) a Group 4 amino acid residue mutated to Group6 amino acid residue at position 153;
 - (m) a Group 4 amino acid residue mutated to a
 Group 5 amino acid residue at position 153;
 - (n) a Group 4 amino acid residue mutated to a Group 1 amino acid residue at position 281;
 - (o) a Group 3 amino acid residue mutated to a Group 2 amino acid residue at position 367;

- 5 (p) a Group 3 amino acid residue mutated to a Group 6 amino acid residue at position 367;
 - (q) a Group 1 amino acid residue mutated to Group 4 amino acid residue at position 389; and
 - (r) a Group 1 amino acid residue mutated to a Group
- 10 2 amino acid residue at position 389.
- 40. The nucleic acid of claim 39, wherein the polynucleotide encodes a variant protein of the lovE protein having a Group 6 amino acid residue mutated to a Group 2 amino acid residue at position 31.
 - 41. The nucleic acid of claim 40 having the mutation F31L.
- 20 42. The nucleic acid of claim 39, wherein the polynucleotide encodes a variant protein of the lovE protein having a Group 3 amino acid residue mutated to a Group 5 amino acid residue at position 41.
- 25 43. The nucleic acid of claim 42 having the mutation Q41K or Q41R.
- 44. The nucleic acid of claim 39, wherein the polynucleotide encodes a variant protein of the lovE protein having a Group 4 amino acid residue mutated to a Group 2 amino acid residue at position 52.
 - 45. The nucleic acid of claim 44 having the mutation T52I.
 - 46. The nucleic acid of claim 39, wherein the polynucleotide encodes a variant protein of the lovE protein having a Group 4 amino acid residue mutated to a Group 3 amino acid residue at position 52.
 - 47. The nucleic acid of claim 46 having the mutation T52N.

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